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09/618,129	07/17/2000	Xiao Bing Wang	TRIMI	8510
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THOMAS, KAYDEN, HORSTEMEYER & RISLEY, LLP 100 GALLERIA PARKWAY, NW			SPIEGLER, ALEXANDER H	
STE 1750 ATLANTA, GA 30339-5948		ART UNIT	PAPER NUMBER	
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Please find below and/or attached an Office communication concerning this application or proceeding.

	Application No.	Applicant(s)	
	09/618,129	WANG, XIAO BING	
Office Action Summary	Examiner	Art Unit	
	Alexander H. Spiegler	1637	
The MAILING DATE of this commun Period for Reply	nication appears on the cover sheet with	the correspondence address	
A SHORTENED STATUTORY PERIOD F THE MAILING DATE OF THIS COMMUN Extensions of time may be available under the provisions after SIX (6) MONTHS from the mailing date of this commodified the period for reply specified above is less than thirty (3) If NO period for reply is specified above, the maximum is a Failure to reply within the set or extended period for reply Any reply received by the Office later than three months earned patent term adjustment. See 37 CFR 1.704(b).	IICATION. s of 37 CFR 1.136(a). In no event, however, may a repl munication. 30) days, a reply within the statutory minimum of thirty (; tatutory period will apply and will expire SIX (6) MONTH y will. by statute, cause the application to become ARAN	ly be timely filed (30) days will be considered timely. 185 from the mailing date of this communication.	
Status	· ·		
1) Responsive to communication(s) file	ed on 26 November 2003.		
	2b)⊠ This action is non-final.		
3) Since this application is in condition	for allowance except for formal matters	s, prosecution as to the merits is	
closed in accordance with the practi	ice under <i>Ex par</i> te <i>Quayl</i> e, 1935 C.D. 1	11, 453 O.G. 213.	
Disposition of Claims			
4)	re withdrawn from consideration. is/are rejected.		
Application Papers			
9)⊠ The specification is objected to by the			
10)☐ The drawing(s) filed on is/are:			
	ction to the drawing(s) be held in abeyance.		
	the correction is required if the drawing(s)		
11)☐ The oath or declaration is objected to	by the Examiner. Note the attached O	rffice Action or form PTO-152.	
Priority under 35 U.S.C. § 119			
2. Certified copies of the priority3. Copies of the certified copies of application from the Internation	documents have been received. documents have been received in Appl of the priority documents have been rec nal Bureau (PCT Rule 17.2(a)).	lication No ceived in this National Stage	
* See the attached detailed Office action	n for a list of the certified copies not rec	eived.	
Attachment(s)		•	
1) Notice of References Cited (PTO-892)	4) 🔲 Interview Sumr	mary (PTO-413)	
 Notice of Draftsperson's Patent Drawing Review (P²) Information Disclosure Statement(s) (PTO-1449 or Paper No(s)/Mail Date <u>5/28/03</u>. 		lail Date mal Patent Application (PTO-152)	

DETAILED ACTION

Continued Examination Under 37 CFR 1.114

1. A request for continued examination under 37 CFR 1.114, including the fee set forth in 37 CFR 1.17(e), was filed in this application after final rejection. Since this application is eligible for continued examination under 37 CFR 1.114, and the fee set forth in 37 CFR 1.17(e) has been timely paid, the finality of the previous Office action has been withdrawn pursuant to 37 CFR 1.114. Applicant's submission filed on November 26, 2003 has been entered.

Status of the Application

2. Currently, claims 2-3, 5, 8-9, 11-37 and 39-41 are pending and are rejected herein.

Information Disclosure Statement

3. The information disclosure statement filed on May 28, 2003 complies with CFR 1.97, 1.98, and M.P.E.P. 609, and has been considered (see enclosed signed PTO-1449).

Specification

- 4. The disclosure is objected to because of the following informalities:
- A) The use of trademarks have been noted in this application. (See, for example, pages 19 "QIAGEN", "PERKIN ELMER", page 20, "PROMEGA", "INVITROGEN", etc.). These should be capitalized wherever it appears and be accompanied by the generic terminology.

Although the use of trademarks is permissible in patent applications, the proprietary nature of the marks should be respected and every effort made to prevent their use in any manner which might adversely affect their validity as trademarks.

B) Claim 15 recites, "on or more", which should recite, "one or more".

Claim Rejections - 35 USC § 112

- 5. The following is a quotation of the second paragraph of 35 U.S.C. 112:
 - The specification shall conclude with one or more claims particularly pointing out and distinctly claiming the subject matter which the applicant regards as his invention.
- 6. Claims 2-3, 5, 8-9 and 11-37 are rejected under 35 U.S.C. 112, second paragraph, as being indefinite for failing to particularly point out and distinctly claim the subject matter which applicant regards as the invention.
- A) Claims 2-3, 5, 8-9 and 11-37 over "the variant target nucleotide base" because this recitation lacks antecedent basis.
- B) Claim 13 over "the 'Klenow fragment' thereof" because this recitation lacks antecedent basis.
- C) Claims 28, 32, 34 and 36 over "extragenomic" because it is not clear as to what this recitation means, there is no definition of this term in the specification, and it is not an art recognized term. Accordingly, "extragenomic" is indefinite.

Claim Rejections - 35 USC § 102

7. The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis for the rejections under this section made in this Office action:

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(b) the invention was patented or described in a printed publication in this or a foreign country or in public use or on sale in this country, more than one year prior to the date of application for patent in the United States.

8. Claims 2-3, 5, 8-9, 11-37 and 39-41 are rejected under 35 U.S.C. 102(e) as being anticipated by Soderlund (US 6,013,431).

Regarding Claims 2-3, 5, 37 and 39-41, Soderland teaches a method for detecting the presence of a nucleic acid of interest having a variant of a known nucleotide base in a predetermined position of a known nucleic acid comprising:

- (a) obtaining a nucleic acid of interest having a target nucleotide base at the predetermined position in a template of the nucleic acid of interest;
- (b) preparing an unlabeled primer complementary to a sequence immediately upstream of the target nucleotide base;
- (c) treating a sample containing the nucleic acid of interest, if the nucleic acid is double-stranded, so as to obtain unpaired nucleotide bases spanning the predetermined position, or directly employing step (d) if the nucleic acid of interest is single-stranded;
- (d) annealing the primer from (b) with the nucleic acid of interest from (c) to obtain a primer-nucleic acid duplex, wherein the target nucleotide base in the nucleic acid of interest is the first unpaired base immediately downstream of the 3' end of the primer;
- (e) mixing the primer-nucleic acid duplex from (d) with a primer extension reaction reagent comprising: (i) three types of non-terminator nucleotides that are not complementarily matched to the known nucleotide base in the predetermined position of the nucleic acid of interest, wherein at least one type of the non-terminator nucleotide is labeled with a detectable marker; and optionally (ii) one type of terminator nucleotide that is complementarily matched to the known nucleotide base in the predetermined

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position of the nucleic acid of interest, wherein the terminator nucleotide is not labeled;

- (f) extending the primer extension reaction by enzymatic or chemical means to form a labeled primer extension product comprising a plurality of labeled non-terminator nucleotides, wherein a labeled primer extension product does not form when the target nucleotide base in the predetermined position of the nucleic acid of interest is the same as the known target nucleotide base in the predetermined position of the known nucleic acid; and
- (g) determining the presence of the nucleic acid of interest having the variant target nucleotide base at the predetermined position in the nucleic acid of interest by detecting the presence of the labeled primer extension product, wherein detecting the labeled primer extension product is not based on size.

(See col. 3, lines 34-58; col. 4, lines 59-63; cols. 7-8, Examples 1-7, and col. 18, ln. 19-53, for example).

With respect to the recitations of "variant" and "known", there are no specific definitions of these recitations in the specification, and accordingly, they have been given the broadest reasonable interpretation consistent with the supporting disclosure and those skilled in the art.

See MPEP 2111. Specifically, the specification states:

By "standard nucleotide base", it includes any known base, which may include wild-type or a known mutant base so long as the base is known and it is desired to know its variant. Thus, as an example, normal base can be a known wild-type base for which a mutation is sought at the position. Inversely, the known base can be a known mutant for which the presence of a wild-type base is sought at the position. Alternatively, the known normal base can be a known mutant for which another mutant variant base is sought. Therefore, the method of the invention can be applied to any known sequence that can be used to determine the presence of any other base variant at the site.

(emphasis added) (page 10-11).

Accordingly, the instant invention is drawn to detecting variations of "known" bases,

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wherein the known base *may* be either a "wild-type" or "mutant" base. Soderlund teaches "a method to determine *a* specific nucleotide *variation in a previously defined region*" (col. 4, lines. 59-60). Thus, Soderlund teaches detecting variations of "known" bases, since Soderlund teaches detecting variations from "previously defined regions". A "previously defined region" (e.g., a previously defined base) is interpreted as a "known" base.

The claims are drawn to detecting the presence of the nucleic acid of interest having the variant target nucleotide base at the predetermined position in the nucleic acid of interest by detecting the presence of the labeled primer extension product. That is, the claims are drawn to detecting a variant of a "known" base by detecting the presence of a labeled extension product. Thus, a labeled primer extension product could be formed by the incorporation of a labeled dNTP, wherein the dNTP corresponds to a variant of a "known" base. Soderlund teaches this detection of a labeled primer extension product, which is indicative of a variant of a "known" base (see, for example, col. 8, lines 49-52).

Regarding Claim 8, Soderlund teaches that two or more differently labeled dNTPs (non-terminator nucleotides) can be added to the primer-nucleic acid duplex, wherein the detection is better interpreted by adding dNTPs that are different than the terminator nucleotide (See col. 8, ln. 58-64, for example).

Regarding Claim 9, Soderlund teaches the use of this invention with various labels such as radioactive or fluorescent labels (see examples 1-7, col. 9-18, for example).

Regarding Claims 11-14, Soderlund teaches that the primer extension reaction can be performed by enzymatic means using template dependent enzymes (i.e. T7 DNA polymerase, T4 DNA polymerase, reverse transcriptase, etc.) (col. 8, ln. 10-17, for example).

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Regarding claims 15-22, Soderlund teaches that the primer may contain an attachment moiety (i.e. biotin, antigens, etc.) (See col. 6, ln. 16-31, for example), that permits affinity separation of the primer from the unincorporated reagent and/or the nucleic acid of interest (col. 6, ln. 53 to col. 7, ln. 26, for example), and furthermore, that a solid support may be used in the separation process (col. 6-7, for example).

Regarding Claims 23-36, Soderlund teaches the source of the target nucleic acid of interest can be any form of RNA or DNA obtained via amplification (for example), from any source such as from a human, animal, plant, or microbe (See cols. 1-5 and Examples 1-7).

Applicant's Arguments

Applicant argues:

- 1) Soderlund is directed to a sequencing method, whereas the instant method is not a sequencing method.
- 2) Soderlund fails to teach a reaction mixture having three types of non-terminator nucleotides that are not complementarily matched to the known nucleotide base. Soderlund teaches "using a primer extension reaction mixture containing a labeled known nucleotide so that the detection of a labeled primer extension product means that target nucleotide at the predetermined position is complementary to the labeled known nucleotide". See Applicant's arguments.
- 3) Soderlund fails to disclose using unlabeled ddNTPs that are complementary to a known nucleotide at a predetermined position so that labeled primer extension products will only form when there is a variation at the predetermined position.

4) Soderlund fails to disclose a primer extension reaction is not formed because Soderlund is directed to determining the identity of the nucleotide at the predetermined position.

Response to Applicants Arguments

Applicant's arguments have been considered, but are not persuasive for the following reasons:

- 1) Both the instant claims and the teachings of Soderlund are drawn to methods of detecting variations of "known" nucleotide bases. Specifically, for example, Claim 37 is drawn to detecting the presence of a nucleic acid interest having a variant of a known nucleotide base in a predetermined position of a known nucleic acid. Soderlund teaches this detection of a variant of a "known" base (see, for example, col. 8, lines 49-52).
- 2) Soderlund does teach the use of three non-terminator nucleotides that are not complementary to the known nucleotide base. See, for example, col. 8, lines 49-52 and lines 58-60. Specifically, the non-terminator nucleotides correspond to a variant of a "known" nucleotide. Applicant's argument with respect to Soderlund's reaction mixture containing a labeled "known" nucleotide is somewhat confusing, since Soderlund is adding a labeled non-terminator (e.g., a labeled dNTP), which is the same type of labeled non-terminator as in the instant invention (e.g., a labeled dNTP, see claim 5). That is, it is assumed that all dNTPs are "known" in the sense that they are dATPs, dTTPs, dGTPs, dCTPs or dUTPs.
- 3) Applicant is arguing limitations not required by the claims, since the claims do not require the use of an unlabeled ddNTPs. At most, Claims 37 and 41 suggest that the use of terminators (e.g., ddNTPs) is "optional", and Claims 39 and 40 do not mention ddNTPs at all. It is also noted the use of term "known" is relative and is not defined in the specification.

Furthermore, as required by the claims, Soderlund teaches the detection of a labeled extension product, wherein the presence of the extension product is indicative of the presence of a variant of a "known" nucleotide base.

4) Like the instant invention, Soderlund's labeled primer extension product will not form if the target nucleotide base in the predetermined position is the same as the known target nucleotide base. For example, Soderlund teaches a labeled extension product when there is a variation of a "known" nucleotide base, and therefore, there will not be a labeled extension product if the target nucleotide base in the predetermined position is the same as the "known" target nucleotide base.

For these reasons, and those of record, the rejection is maintained.

9. Claims 2-9, 11-15, 17, 23-37 and 39-41 are rejected under 35 U.S.C. 102(b) as being anticipated by Fahy et al. (WO 96/30545, previously cited).

Regarding Claims 2-3, 5, 37 and 39-41, Fahy teaches a method for detecting the presence of a nucleic acid of interest having a variant of a known nucleotide base in a predetermined position of a known nucleic acid comprising:

- (a) obtaining a nucleic acid of interest having a target nucleotide base at the predetermined position in a template of the nucleic acid of interest;
- (b) preparing an unlabeled primer complementary to a sequence immediately upstream of the target nucleotide base;
- (c) treating a sample containing the nucleic acid of interest, if the nucleic acid is double-stranded, so as to obtain unpaired nucleotide bases spanning the predetermined position, or directly employing step (d) if the nucleic acid of interest is single-stranded;

(d) annealing the primer from (b) with the nucleic acid of interest from (c) to obtain a primer-nucleic acid duplex, wherein the target nucleotide base in the nucleic acid of interest is the first unpaired base immediately downstream of the 3' end of the primer;

- (e) mixing the primer-nucleic acid duplex from (d) with a primer extension reaction reagent comprising: (i) three types of non-terminator nucleotides that are not complementarily matched to the known nucleotide base in the predetermined position of the nucleic acid of interest, wherein at least one type of the non-terminator nucleotide is labeled with a detectable marker; and optionally (ii) one type of terminator nucleotide that is complementarily matched to the known nucleotide base in the predetermined position of the nucleic acid of interest, wherein the terminator nucleotide is not labeled;
- (f) extending the primer extension reaction by enzymatic or chemical means to form a labeled primer extension product comprising a plurality of labeled non-terminator nucleotides, wherein a labeled primer extension product does not form when the target nucleotide base in the predetermined position of the nucleic acid of interest is the same as the known target nucleotide base in the predetermined position of the known nucleic acid; and
- (g) determining the presence of the nucleic acid of interest having the variant target nucleotide base at the predetermined position in the nucleic acid of interest by detecting the presence of the labeled primer extension product, wherein detecting the labeled primer extension product is not based on size.

(See page 6, lines 31-36, pages 7-8, page 16, lines 17-37, pages 18-19, page 21, lines 27-32, for example).

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With respect to the recitations of "variant" and "known", there are no specific definitions of these recitations in the specification, and accordingly, they have been given the broadest reasonable interpretation consistent with the supporting disclosure and those skilled in the art.

See MPEP 2111. Specifically, the specification states:

By "standard nucleotide base", it includes any known base, which may include wild-type or a known mutant base so long as the base is known and it is desired to know its variant. Thus, as an example, normal base can be a known wild-type base for which a mutation is sought at the position. Inversely, the known base can be a known mutant for which the presence of a wild-type base is sought at the position. Alternatively, the known normal base can be a known mutant for which another mutant variant base is sought. Therefore, the method of the invention can be applied to any known sequence that can be used to determine the presence of any other base variant at the site.

(emphasis added) (page 10-11).

Accordingly, the instant invention is drawn to detecting variations of "known" bases, wherein the known base *may* be either a "wild-type" or "mutant" base. Fahy teaches the dNTPs are not complementary to a "known" target nucleotide base (see page 19, lines 1-17; see also page 16, lines 17-20 and page 21, lines 27-32).

It is noted that, the terms "wild-type" and "mutant" are relative terms, which can be population specific. That is, these terms are not fixed terms; in one population a specific nucleotide may be considered "wild-type", but in a different population the specific nucleotide may be considered "mutant".

It is also noted that Fahy teaches the use of labeled dNTPs for detection of extension products (which is identical to the instant invention), and therefore, the detection the labeled primer extension product is not based on size, but on whether or not the product contains the label. See, for example, page 21, lines 27-32.

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Regarding Claim 8, Fahy teaches the labeled dNTPs (non-terminator nucleotides) are labeled with the same or different detectable markers (See page 21, lines 15-23 and 27-32, for example).

Regarding Claim 9, Fahy teaches the use of this invention with various labels such as radioactive or fluorescent labels (See pages 15-16 and page 21, lines 15-23, for example).

Regarding Claims 11-14, Fahy teaches that the primer extension reaction can be performed by enzymatic means using template dependent enzymes (i.e., T7 DNA polymerase, Klenow fragment, reverse transcriptase, etc.) (See pages 19-20, for example).

Regarding claims 15 and 17, Fahy teaches that the primer may contain biotin (See page 21, lines 15-26, for example).

Regarding Claims 23-36, Fahy teaches the source of the target nucleic acid of interest can be any form of RNA or DNA obtained via amplification (for example), from any source such as from a human, animal, or microbe (See pages 9, 13-15, 17 and 58, for example).

Claim Rejections - 35 USC § 103

- 10. The following is a quotation of 35 U.S.C. 103(a) which forms the basis for all obviousness rejections set forth in this Office action:
 - (a) A patent may not be obtained though the invention is not identically disclosed or described as set forth in section 102 of this title, if the differences between the subject matter sought to be patented and the prior art are such that the subject matter as a whole would have been obvious at the time the invention was made to a person having ordinary skill in the art to which said subject matter pertains. Patentability shall not be negatived by the manner in which the invention was made.
- 11. The factual inquiries set forth in *Graham* v. *John Deere Co.*, 383 U.S. 1, 148 USPQ 459 (1966), that are applied for establishing a background for determining obviousness under 35 U.S.C. 103(a) are summarized as follows:

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- 1. Determining the scope and contents of the prior art.
- 2. Ascertaining the differences between the prior art and the claims at issue.
- 3. Resolving the level of ordinary skill in the pertinent art.
- 4. Considering objective evidence present in the application indicating obviousness or nonobviousness.
- 12. Claims 16 and 18-22 are rejected under 35 U.S.C. 103(a) as being unpatentable over Fahy et al. (WO 96/30545, previously cited), as applied to claims 2-9, 11-15, 17, 23-37 and 39-41 above, and further in view of Soderlund (US 6,013,431).

The teachings of Fahy are presented above. Specifically, Fahy teaches the method of detection using a primer labeled with biotin. Fahy does not teach the primer permits affinity separation of the primer from the unincorporated reagent and/or the nucleic acid of interest or that a solid support may be used in the separation process.

However, Soderlund teaches a detection method wherein the primer may contain an attachment moiety (i.e. biotin, antigens, etc.) (See col. 6, ln. 16-31, for example), that permits affinity separation of the primer from the unincorporated reagent and/or the nucleic acid of interest (col. 6, ln. 53 to col. 7, ln. 26, for example), and furthermore, that a solid support may be used in the separation process (col. 6-7, for example). Soderlund teaches using attachment moieties (which can be used for linkage to a solid support) is advantageous in aiding the detection process by determining which strand the variable nucleotide occurs (col. 6, lines 7-9), purifying the reaction to ensure only bound material is analyzed (col. 6, lines 55-63), and making it possible to reuse the target nucleic acid if multiple determinations are to be performed on the same target sequence of interest (col. 6, lines 63-67).

Accordingly, in view of the teachings of Soderlund, it would have been obvious to one of ordinary skill in the art at the time the invention was made to have modified the method of Fahy

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so as to have used a primer comprising an attachment moiety which permits affinity separation of the primer from the unincorporated reagent and/or the nucleic acid of interest and to have used a solid support in the separation process. One of ordinary skill in the art would have been motivated to modify the method of Fahy in order to have achieved the benefit of aiding the detection process by determining which strand the variable nucleotide occurs, purifying the reaction to ensure only bound material is analyzed, and making it possible to reuse the target nucleic acid if multiple determinations are to be performed on the same target sequence of interest.

Conclusion

13. No claims are allowable.

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Correspondence

Any inquiry concerning this communication or earlier communications from the

examiner should be directed to Alexander H. Spiegler whose telephone number is (571) 272-

0788. The examiner can normally be reached on Monday through Friday, 7:00 AM to 3:30 PM.

If attempts to reach the examiner are unsuccessful, the primary examiner in charge of the

prosecution of this case, Carla Myers, can be reached at (571) 272-0747. If attempts to reach

Carla Myers are unsuccessful, the examiner's supervisor, Gary Benzion can be reached at (571)

272-0782.

Papers related to this application may be faxed to Group 1637 via the PTO Fax Center

using the fax number (703) 872-9306.

Information regarding the status of an application may be obtained from the Patent

Application Information Retrieval (PAIR) system. Status information for published applications

may be obtained from either Private PAIR or Public PAIR. Status information for unpublished

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system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR

system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free).

Alexander H. Spiegler

March 3, 2004

le May 313104

DUBARY EXAMINER